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Total Errors: 0

No. of SeqIDs Defined: 16

Actual SeqID Count: 16

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SCHUIJFFEL, Danielle Francisca

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		aca gat gtt tct gca at Thr Asp Val Ser Ala Il 205	
		aaa gcg gaa gct cca aa Lys Ala Glu Ala Pro Ly 220	
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305 310 315 320

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420 425 430

Val Lys Asp Leu Ala Thr Arg Ser Arg Asp Arg Lys Ile Lys Ala Asp 435 440 445 Glu Met Glu Gly Ser Thr Phe Thr Val Ser Asn Leu Gly Ala Tyr Gly 450 455 460 Val Glu Ser Phe Thr Ser Ile Ile Asn Gln Pro Asn Ser Cys Ile Leu 465 470 475 480 Ser Val Gly Ala Ile Val Glu Lys Pro Val Val Lys Asn Gly Gln Ile 485 490 495 Val Val Gly His Thr Met Lys Leu Cys Leu Ala Cys Asp His Arg Thr 505 500 510 Val Asp Gly Ala Thr Gly Ser Thr Phe Leu Gln Thr Leu Lys Gln Tyr 515 520 525 Leu Glu Thr Pro Met Ser Met Leu Val 530 535 <210> 3 <211> 1572 <212> DNA <213> Ornithobacterium rhinotracheale <220> <221> CDS <222> (1)..(1572) <400> 3 atg aaa ata aat tac aaa aat ata ctt tta agt gct agc gtt ctc ttt 48 Met Lys Ile Asn Tyr Lys Asn Ile Leu Leu Ser Ala Ser Val Leu Phe 10 15 1 ttt gca gca tgt agc gat ttt gat tac aat gta gaa aac cca aac ctc 96 Phe Ala Ala Cys Ser Asp Phe Asp Tyr Asn Val Glu Asn Pro Asn Leu 25 20 30 acg aag gga gag gct gat ttc tct aaa tat gta gct tta gga aat tct 144Thr Lys Gly Glu Ala Asp Phe Ser Lys Tyr Val Ala Leu Gly Asn Ser 35 40 45 ctc act tct ggt tat tca gac gga gcc tta tat cgc tcg gca caa gag 192 Leu Thr Ser Gly Tyr Ser Asp Gly Ala Leu Tyr Arg Ser Ala Gln Glu

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55

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